



-14 -10
AC Leu Ala Leu Cys Leu Leu Thr Phe Thr Ser Ser Ala Thr Ala 44
CUG GCG CUG UGC CUG CUC ACC UUC ACC AGC UCU GCC ACG GCU

1 10
Gly Pro Glu Thr Leu Cys Gly Ala Glu Leu Val Asp Ala Leu Gln 89
GGA CCG GAG ACG CUC UGC GGG GCU GAG CUG GUG GAU GCU CUU CAG

20 30
Phe Val Cys Gly Asp Arg Gly Phe Tyr Phe Asn Lys Pro Thr Gly 134
UUC GUG UGU GGA GAC AGG GGC UUU UAU UUC AAC AAG CCC ACA GGG

40
Tyr Gly Ser Ser Ser Arg Arg Ala Pro Gln Thr Gly Ile Val Asp 179
UAU GGC UCC AGC AGU CGG AGG GCG CCU CAG ACA GGU AUC GUG GAU

50 60
Glu Cys Cys Phe Arg Ser Cys Asp Leu Arg Arg Leu Glu Met Tyr 224
GAG UGC UGC UUC CGG AGC UGU GAU CUA AGG AGG CUG GAG AUG UAU

70
Cys Ala Pro Leu Lys Pro Ala Lys Ser Ala Arg Ser Val Arg Ala 269
UGC GCA CCC CUC AAG CGU GCG AAG UCA GCU CCG UCU GUC CGU GCC

80 90
Gln Arg His Thr Asp Met Pro Lys Thr Gln Lys Glu Val His Leu 314
CAG CCG CAC ACC GAG AUG CCG AAG ACC CAG AAG GAA GUA CAU UUG

100 105
Lys Asn Ala Ser Arg Gly Ser Ala Gly Asn Lys Asn Tyr Arg Met 359
AAG AAC GCA AGU AGA GGG AGU GCA GGA AAC AAG AAC UAC AGG AUG

AM
UAG GAAGACCCUCCUGAGGAGUGAAGAGUGACAUGCCACCGCA GGAUCCUUUGCUCUGCA 419
CGAGUUACCGUUAUUUUUGGAA CACCUACCAAAAAUAAGUUUGAUACAUAUUAAAAAG 479
AUGGGCGUUUCGCCCAUUGAAAUACACAA GUAAACAUUCCAACAUUGUCUUUAGGAGUGA 539
UUUGCACCUUGCAAAAAUGGUCCUGGAGUUGGUGAGAUUG CUGUUGAUCUUUAUCAAUA 599
UGUUCUAUAAAAAAAAAAAAAA

FIG. 1.

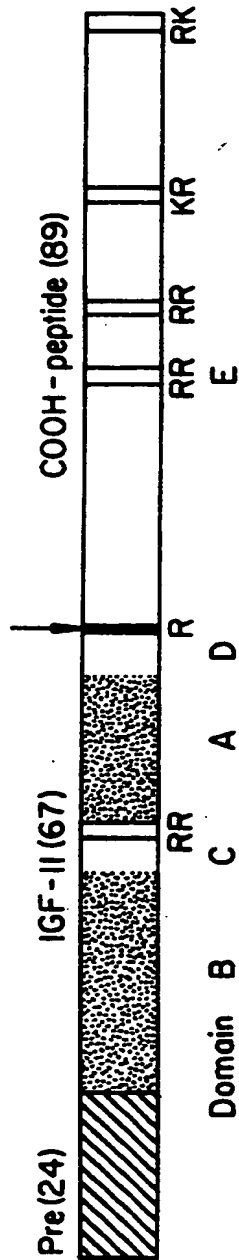


FIG.—3.

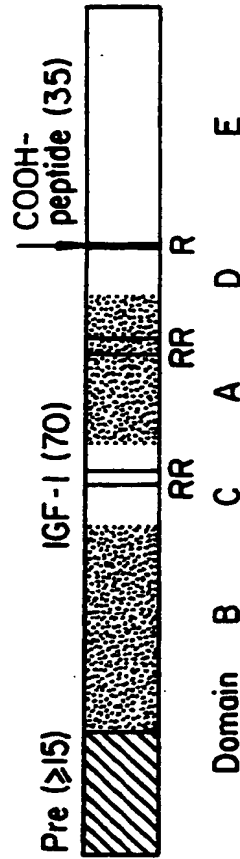


FIG.—4.